

IN THE CLAIMS

Please amend the claims as follows:

Claims 1-29(Canceled)

Claim 30 (Currently Amended): A process for producing an L-amino acid[[s]] comprising:
culturing a bacterial cell in a medium suitable for producing an L-amino acid[[s]], and recovering or isolating said L-amino acid,
wherein said bacterial cell ~~comprises an attenuated *ccpA2* gene~~ has been modified to eliminate or reduce the expression of the CcpA2 polypeptide compared to the corresponding unmodified bacterium, and
wherein said CcpA2 polypeptide exhibits catabolite control activity and is encoded by a polynucleotide which is at least 95% identical to SEQ ID NO: 1.

Claim 31 (Currently Amended): The process of Claim 30, wherein said bacterial cell is a ~~*Coryneform*~~ *coryneform* bacterium ~~or *Brevibacterium*~~.

Claim 32 (Currently Amended): The process of Claim 31, wherein said bacterial cell is selected from the group consisting of ~~*Coryneform*~~ *Corynebacterium glutamicum*, *Corynebacterium acetoglutamicum*, *Corynebacterium acetoacidophilum*, *Corynebacterium melassecola*, *Corynebacterium thermoaminogenes*, *Brevibacterium flavum*, *Brevibacterium lactofermentum*, and *Brevibacterium divaricatum*.

Claim 33 (Currently Amended): The process of Claim 30, wherein the CcpA2 polypeptide comprises SEQ ID NO: 2 ~~said *ccpA2* gene comprises the polynucleotide sequence of SEQ ID NO: 1.~~

Claim 34 (Currently Amended): The process of Claim 30, wherein ~~said~~ the L-amino acid is L-lysine.

Claims 35- 37(Canceled)

Claim 38 (New): The process of Claim 30, wherein the L-amino acid is not L-lysine.

Claim 39 (New): The process of Claim 30, wherein the expression of the CcpA2 polypeptide has been eliminated.

Claim 40 (New): The process of Claim 30, wherein the expression of the CcpA2 polypeptide has been attenuated.

Claim 41 (New): The process of Claim 30, wherein the expression of the CcpA2 polypeptide has been attenuated by genetic modification of a signal structure involved in expression of the CcpA2 coding sequence.

Claim 42 (New): The process of Claim 30, wherein the expression of the CcpA2 polypeptide has been eliminated or attenuated by modification of the CcpA2 coding sequence.

Claim 43 (New): The process of Claim 30, wherein said bacterium contains one or more coding sequences from *lysC*, *dapA*, *eno*, *zwf*, *dapD*, *dapE*, *gap*, *pyc*, *mgo*, *zal*, or *lysE*, wherein said one or more coding sequences is over-expressed compared to a corresponding unmodified bacterium.

Claim 44 (New): The process of Claim 30, wherein said bacterium has had one or more coding sequences from *pck*, *pgi*, *poxB*, or *zwa2* eliminated or attenuated compared to a corresponding unmodified strain.

Claim 45 (New): The process of Claim 30, wherein said L-amino acid is produced by a batch fermentation process.

Claim 46 (New): The process of Claim 30, wherein said L-amino acid is produced by a fed batch fermentation process.

Claim 47 (New): The process of Claim 30, wherein said L-amino acid is produced by a repeated fed fermentation process.

Claim 48 (New): The process of Claim 30, wherein said L-amino acid is recovered from the culture medium.

Claim 49 (New): The process of Claim 30, wherein said L-amino acid is recovered along with at least a portion of the biomass of the culture medium.

Claim 50 (New): A process for producing an L-amino acid comprising:
culturing a bacterial cell in a medium suitable for producing an L-amino acid, and
recovering or isolating said L-amino acid;
wherein said bacterial cell has been modified to eliminate or reduce the
expression of the CcpA2 polypeptide compared to the corresponding unmodified bacterium,
and

wherein said CcpA2 polypeptide is encoded by a polynucleotide which may be
obtained from *Brevibacterium* or *Corynebacterium* by PCR using the oligonucleotides of
SEQ ID NOS: 3 and 4.

Claim 51 (New): The process of claim 30, wherein said CcpA2 polypeptide is
encoded by a polynucleotide which comprises nucleotides 241 to 1278 of SEQ ID NO: 1.